

2459-1-003 SeqList 09-15-08.txt SEQUENCE LISTING

<110> Zhou, Ming-Ming Aggarwal, Aneel Methods of Identifying Modulators of Bromodomains <120> <130> 2459-1-003 <140> 09510314 <141> 2008-03-19 <150> 09/510,314 2000-02-22 <151> <160> 44 <170> PatentIn version 3.0 <210> 3014 <211> <212> DNA <213> Homo sapiens <400> 1 60 ggggccgcgt cgacgcggaa aagaggccgt ggggggcctc ccagcgctgg cagacaccgt gaggetggca geegeeggea egeacaceta gteegeagte eegaggaaca tgteegeage 120 cagggcgcgg agcagagtcc cgggcaggag aaccaaggga gggcgtgtgc tgtggcggcg 180 gcggcagcgg cagcggagcc gctagtcccc tccctcctgg gggagcagct gccgccgctg 240 300 ccgccgccgc caccaccatc agcgcgcggg gcccggccag agcgagccgg gcgagcggcg 360 420 cctgacactc ggcgcctcct gccgtgctcc ggggcggcat gtccgaggct ggcggggccg 480 ggccgggcgg ctgcgggca ggagccgggg caggggccgg gcccggggcg ctgccccgc agcctgcggc gcttccgcc gcgccccgc agggctcccc ctgcgccgct gccgccgggg gctcgggcgc ctgcggtccg gcgacggcag tggctgcagc gggcacggcc gaaggaccgg gaggcggtgg ctcggcccga atcgccgtga agaaagcgca actacgctcc gctccgcggg 540 600 660 720 ccaagaaact ggagaaactc ggagtgtact ccgcctgcaa ggccgaggag tcttgtaaat 780 gtaatggctg gaaaaaccct aacccctcac ccactcccc cagagccgac ctgcagcaaa 840 taattotcag tctaacagaa tcctgtcgga gttgtagcca tgccctagct gctcatgttt cccacctgga gaatgtgtca gaggaagaaa tgaacagact cctgggaata gtattggatg 900 960 tggaatatct ctttacctgt gtccacaagg aagaagatgc agataccaaa caagtttatt tctatctatt taagctcttg agaaagtcta ttttacaaag aggaaaacct gtggttgaag gctctttgga aaagaaaccc ccatttgaaa aacctagcat tgaacagggt gtgaataact ttgtgcagta caaatttagt cacctgccag caaaagaaag gcaacaata gttgagttgg caaaaaatgtt cctaaaccgc atcaactatt ggcatctgga ggcaccatct caacgaagac 1020 1080 1140 1200 1260 tgcgatctcc caatgatgat atttctggat acaaagagaa ctacacaagg tggctgtgtt 1320 actgcaacgt gccacagitc tgcgacagtc tacctcggta cgaaaccaca caggtgtttg 1380 ggagaacatt gcttcgctcg gtcttcactg ttatgaggcg acaactcctg gaacaagcaa 1440 1500 gacaggaaaa agataaactg cctcttgaaa aacgaactct aatcctcact catttcccaa āattīctgtc cātgctagaā gaagaagtat atagtcaaaa ctctcccatc tgggatcagg 1560 attitictic agcitagaa gaagaagtat atagicaada ciciccati tyggatcagg attitictic agcitagaa gaagaagtat atagicaada ciciccati tyggatcagg cicctgtggc tyggacaatt tcatacaatt caacctcatc ticccttgag cagccaaacg cagggagcag cagtcitgcc tycaaagcci citctggact tygggcaaac ccaggagaaa agaggaaaat gactgattic catgitictgg aggaggccaa gaaaccccga gitatygggg atatticcgat ggaattaatc aacgaggita tyticaccat cacggaccct gcagcaatgc 1620 1680 1740 1800 1860 ttggaccāga gāccaatttt ctgīcāgcac actcggccag ggaīgaggcg gcaāggttgg 1920 aagagcgcag gggtgtaatt gaatttcacg tggttggcaa ttccctcaac cagaaaccaa acaagaagat cctgatgtgg ctggttggcc tacagaacgt tttctcccac cagctgcccc gaatgccaaa agaatacatc acacggctcg tctttgaccc gaaacacaaa acccttgctt taattaaaga tggccgtgtt attggtggta tctgtttccg tatgttccca tctcaaggat tcacagagat tgtcttctgt gctgtaacct caaatggaca agtcaagggc tatggaacac 1980 2040 2100 2160 2220 acctgatgaa tcatttgaaa gaatatcaca taaagcatga catcctgaac ttcctcacat 2280

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Ser Gly Ala Cys Gly Pro Ala Thr Ala Val Ala Ala Ala Gly Thr Ala 50 55 60
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Gln Leu Arg Ser Ala Pro Arg Ala Lys Lys Leu Glu Lys Leu Gly Val
85 90 95
Tyr Ser Ala Cys Lys Ala Glu Glu Ser Cys Lys Cys Asn Gly Trp Lys 100 105 110
Asn Pro Asn Pro Ser Pro Thr Pro Pro Arg Ala Asp Leu Gln Gln Ile
115 120 125
Ile Val Ser Leu Thr Glu Ser Cys Arg Ser Cys Ser His Ala Leu Ala
130 135 140
Ala His Val Ser His Leu Glu Asn Val Ser Glu Glu Glu Met Asn Arg
145 150 155 160
Leu Leu Gly Ile Val Leu Asp Val Glu Tyr Leu Phe Thr Cys Val His
165 170 175
Lys Glu Glu Asp Ala Asp Thr Lys Gln Val Tyr Phe Tyr Leu Phe Lys
180 185 190
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195 200 205
Ser Leu Glu Lys Lys Pro Pro Phe Glu Lys Pro Ser Ile Glu Gln Gly 210 215 220
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One may be missing.
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One, two, or three may be missing.
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       Xaa represents an acetyl lysine.
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Pro Val Lys Arg Thr Glu Ala Pro Gly Tyr Tyr Glu Val Ile Arg Ser
35 40 45
Pro Met Asp Leu Lys Thr Met Ser Glu Arg Leu Lys Asn Arg Tyr Tyr 50 55 60
Val Ser Lys Lys Leu Phe Met Ala Asp Leu Gln Arg Val Phe Thr Asn 65 70 75 80
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2459-1-003 SeqList 09-15-08.txt

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<213> Homo sapiens

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Leu Leu Ala Gln Ile Lys Ser His Pro Ser Ala Trp Pro Phe Met Glu $20 \hspace{1cm} 25 \hspace{1cm} 30$

Pro Val Lys Lys Ser Glu Ala Pro Asp Tyr Tyr Glu Val Ile Arg Phe 35 40 45

Pro Ile Asp Leu Lys Thr Met Thr Glu Arg Leu Arg Ser Arg Tyr Tyr 50 60

Val Thr Arg Lys Leu Phe Val Ala Asp Leu Gln Arg Val Ile Ala Asn 65 70 75 80

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Val Asn Lys Asp Asp Val Pro Asp Tyr Tyr Asp Val Ile Thr Asp Pro 35 40 45

Asp Lys Asp Gln Phe Ile Lys Asp Val Lys Arg Ile Phe Thr Asn Ala 65 70 75 80

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35 40 45

Met Asp Leu Ser Thr Met Glu Ile Lys Leu Glu Ser Asn Lys Tyr Gln 50 . 55 60

Lys Met Glu Asp Phe Ile Tyr Asp Ala Arg Leu Val Phe Asn Asn Cys 75 80

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Lys Ser Pro Met Asp Leu Ser Thr Ile Lys Arg Lys Leu Asp Thr Gly 50 55 60

Gln Tyr Gln Glu Pro Trp Gln Tyr Val Asp Asp Ile Trp Leu Met Phe 65 70 75 80

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Lys Asn Pro Met Asp Leu Ser Thr Ile Lys Arg Lys Leu Asp Thr Gly 50 60 .

Gln Tyr Gln Glu Pro Trp Gln Tyr Val Asp Asp Val Trp Leu Met Phe 70 75 80

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Cys Ser Lys Leu Ala Glu Val Phe Glu Gln Glu Ile Asp Pro Val Met 100 105 110

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Lys Asn Pro Met Asp Leu Ser Thr Ile Lys Arg Lys Leu Asp Thr Gly 50 55 60

Gln Tyr Gln Glu Pro Trp Gln Tyr Val Asp Asp Val Arg Leu Met Phe 65 70 75 80

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35 40 45
Pro Met Asp Leu Gln Thr Leu Arg Glu Asn Val Arg Lys Arg Leu Tyr 50 60
Pro Ser Arg Glu Glu Phe Arg Glu His Leu Glu Leu Ile Val Lys Asn 65 70 75 80
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2459-1-003 SeqList 09-15-08.txt
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<212> PRT

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His Pro Val Asn Lys Lys Phe Val Pro Asp Tyr Tyr Lys Val Ile Val 35 40 45

Asn Pro Met Asp Leu Glu Thr Ile Arg Lys Asn Ile Ser Lys His Lys 50 55 60

Tyr Gln Ser Arg Glu Ser Phe Leu Asp Asp Val Asn Leu Ile Leu Ala 65 70 75 80

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Gln Glu Ile Val Asn Val Cys Tyr Gln Thr Leu Thr Glu Tyr Asp 100 105 110

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<212> PRT

<213> Mesocricetus auratus

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His Pro Val Asn Lys Lys Phe Val Pro Asp Tyr Tyr Lys Val Ile Val 35 40 45

Ser Pro Met Asp Leu Glu Thr Ile Arg Lys Asn Ile Ser Lys His Lys 50 60 Page 11

Tyr Gln Ser Arg Glu Ser Phe Leu Asp Asp Val Asn Leu Ile Leu Ala 80

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20 25 30

Val Asp Ala Val Lys Leu Gly Leu Pro Asp Tyr His Lys Ile Ile Lys
35 40 45

Gln Pro Met Asp Met Gly Thr Ile Lys Arg Arg Leu Glu Asn Asn Tyr 50 55 60

Tyr Trp Ala Ala Ser Glu Cys Met Gln Asp Phe Asn Thr Met Phe Thr 65 70 75 80

Asn Cys Tyr Ile Tyr Asn Lys Pro Thr Asp Asp Ile Val Leu Met Ala 85 90 95

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Val Asp Ala Ile Lys Leu Asn Leu Pro Asp Tyr His Lys Ile Ile Lys 35 40 45

Asn Pro Met Asp Met Gly Thr Ile Lys Lys Arg Leu Glu Asn Asn Tyr 50 60

Tyr Trp Ser Ala Ser Glu Cys Met Gln Asp Phe Asn Thr Met Phe Thr 65 70 75 80

Asn Cys Tyr Ile Tyr Asn Lys Pro Thr Asp Asp Ile Val Leu Met Ala Page 12 Gln Ala Leu Glu Lys Ile Phe Leu Gln Lys Val Ala Gln Met Pro 100 105 110

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<211> 111

<212> PRT

<213> Drosophila melanogaster

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Gln Pro Met Asp Met Gly Thr Ile Lys Lys Arg Leu Glu Asn Asn Tyr 50 55 60

Tyr Trp Ser Ala Lys Glu Thr Ile Gln Asp Phe Asn Thr Met Phe Asn 65 70 75 80

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Pro Val Lys Leu Asp Ile Pro Phe Tyr Phe Asn Tyr Ile Lys Arg Pro 35 40 45

Met Asp Leu Ser Thr Ile Glu Arg Lys Leu Asn Val Gly Ala Tyr Glu 50 60

Val Pro Glu Gln Ile Thr Glu Asp Phe Asn Leu Met Val Asn Asn Ser 65 70 75 80

Ile Lys Phe Asn Gly Pro Asn Ala Gly Ile Ser Gln Met Ala Arg Asn 85 90 95

Ile Gln Ala Ser Phe Glu Lys His Met Leu Asn Met Pro 100 105

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Ile Lys His Pro Met Asp Leu Ser Thr Val Lys Arg Lys Met Glu Asn 50 60

Arg Asp Tyr Arg Asp Ala Gln Glu Phe Ala Ala Asp Val Arg Leu Met 65 70 75 80

Phe Ser Asn Cys Tyr Lys Tyr Asn Pro Pro Asp His Asp Val Val Ala 85 90 95

Met Ala Arg Lys Leu Gln Asp Val Phe Glu Phe Arg Tyr Ala Lys Met 100 105 110

Pro

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Ile Lys Lys Pro Met Asp Leu Gly Thr Val Lys Arg Lys Met Asp Asn 50 55 60
Arg Glu Tyr Lys Ser Ala Pro Glu Phe Ala Ala Asp Val Arg Leu Ile
65 70 75 80
Phe Thr Asn Cys Tyr Lys Tyr Asn Pro Pro Asp His Asp Val Val Ala
85 90 95
Met Gly Arg Lys Leu Gln Asp Val Phe Glu Met Arg Tyr Ala Asn Ile
100 105 110
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        Saccharomyces cerevisiae
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Lys Glu Leu Met Ala Lys Lys His Ala Ser Tyr Asn Tyr Pro Phe Leu
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Glu Pro Val Asp Pro Val Ser Met Asn Leu Pro Thr Tyr Phe Asp Tyr 35 40 45
Val Lys Glu Pro Met Asp Leu Gly Thr Ile Ala Lys Lys Leu Asn Asp 50 55 60
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Trp Gln Tyr Gln Thr Met Glu Asp Phe Glu Arg Glu Val Arg Leu Val 65 70 75 80

Phe Lys Asn Cys Tyr Thr Phe Asn Pro Asp Gly Thr Ile Val Asn Met $85 \hspace{1cm} 90 \hspace{1cm} 95$

Met Gly His Arg Leu Glu Glu Val Phe Asn Ser Lys Trp Ala Asp Arg 100 105 110

Pro

2459-1-003 SeqList 09-15-08.txt

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Pro Leu Ser Glu Val Pro Asp Tyr Leu Asp His Ile Lys Lys Pro Met 35 40 45

Asp Phe Phe Thr Met Lys Gln Asn Leu Glu Ala Tyr Arg Tyr Leu Asn 50 55 60

Phe Asp Asp Phe Glu Glu Asp Phe Asn Leu Ile Val Ser Asn Cys Leu 65 70 75 80

Lys Tyr Asn Ala Lys Asp Thr Ile Phe Tyr Arg Ala Ala Val Arg Leu 85 90 95

Arg Glu Gln Gly Gly Ala Val Val Arg Gln Ala Arg 100 105

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<400> 28

Ser Glu Asp Gln Glu Ala Ile Gln Ala Gln Lys Ile Trp Lys Lys Ala 1 10 15

Ile Met Leu Val Trp Arg Ala Ala Ala Asn His Arg Tyr Ala Asn Val 20 25 30

Phe Leu Gln Pro Val Thr Asp Asp Ile Ala Pro Gly Tyr His Ser Ile 35 40 45

Val Gln Arg Pro Met Asp Leu Ser Thr Ile Lys Lys Asn Ile Glu Asn 50 60

Gly Leu Ile Arg Ser Thr Ala Glu Phe Gln Arg Asp Ile Met Leu Met 65 70 75 80

Phe Gln Asn Ala Val Met Tyr Asn Ser Ser Asp His Asp Val Tyr His 85 90 95

Met Ala Val Glu Met Gln Arg Asp Val Leu Glu Gln Ile Gln Gln Phe 100 105 110

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2459-1-003 SeqList 09-15-08.txt

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Phe Ile Arg Ala Pro Lys Arg Arg Asn Gln Pro Asp Tyr Tyr Glu Val 35 40 45

Val Ser Gln Pro Ile Asp Leu Met Lys Ile Gln Gln Lys Leu Lys Met 50 55 60

Glu Glu Tyr Asp Asp Val Asn Val Leu Thr Ala Asp Phe Gln Leu Leu 65 70 75 80

Phe Asn Asn Ala Lys Ala Tyr Tyr Lys Pro Asp Ser Pro Glu Tyr Lys 85 90 95

Ala Ala Cys Lys Leu Trp Glu Leu Tyr Leu 100 105

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Gln Lys Leu Pro Ser Lys Val Gln Tyr Pro Asp Tyr Tyr Ala Ile Ile 35 40 45

Lys Glu Pro Ile Asp Leu Lys Thr Ile Ala Gln Arg Ile Gln Asn Gly 50 60

Thr Tyr Lys Ser Ile His Ala Met Ala Lys Asp Ile Asp Leu Leu Ala 65 70 75 80

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Ala Asn Ala Ile Lys Lys Ile Phe Asn Met Lys Lys Ala Glu Ile Glu 100 105 110

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20 25 30
Phe Gln Leu Pro Ser Lys Lys Lys Tyr Pro Asp Tyr Tyr Gln Gln Ile 35 40 45
Lys Thr Pro Ile Ser Leu Gln Gln Ile Arg Ala Lys Leu Lys Asn His 50 60
Glu Tyr Glu Thr Leu Asp Gln Leu Glu Ala Asp Leu Asn Leu Met Phe 65 70 75 80
Glu Asn Ala Lys Arg Tyr Asn Val Pro Asn Ser Ala Ile Tyr Lys Arg
85 90 95
Val Leu Lys Met Gln Gln Val Met Gln Ala Lys Lys Lys Glu Leu Ala
100 105 110
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113
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20 25 30
Phe Met Val Lys Pro Ser Lys Lys Asp Tyr Pro Asp Tyr Tyr Lys Ile 35 \hspace{1cm} 40 \hspace{1cm} 45
Ile Leu Glu Pro Met Asp Leu Lys Met Ile Glu His Asn Ile Arg Asn 50 55 60
Asp Lys Tyr Val Gly Glu Glu Ala Met Ile Asp Asp Met Lys Leu Met 65 70 75 80
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Asp Ala His Met Leu Glu Lys Ile Leu Lys Glu Lys Arg Lys Glu Leu
100 105 110
Gly
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        33
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        115
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1 5 10
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Ile Thr Ile Lys Lys Pro Val Asp Met Glu Lys Ile Arg Ser His Met 60

Met Ala Asn Lys Tyr Gln Asp Ile Asp Ser Met Val Glu Asp Phe Val 75

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Gly Glu Týr Thr Thr Leu Glu Ser Phe Val Lys Asp Leu Asn Gln Met 65 70 75 80
Phe Ile Asn Ala Lys Thr Tyr Asn Ala Pro Gly Ser Phe Val Tyr Glu
85 90 95
Asp Ala Glu Lys Leu Ser Gln Leu Ser Ser Ser Leu Ile Ser Ser Phe 100 \hspace{1cm} 105 \hspace{1cm} 110
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Phe Ile Asp Leu Pro Ser Lys Arg Leu Tyr Pro Asp Tyr Tyr Glu Ile 35 40 45
Ile Lys Ser Pro Met Thr Ile Lys Met Leu Glu Lys Arg Phe Lys Lys 50 55 60
Gly Glu Tyr Thr Thr Leu Glu Ser Phe Val Lys Asp Leu Asn Gln Met 70 75 80
Phe Ile Asn Ala Lys Thr Tyr Asn Ala Pro Gly Ser Phe Val Tyr Glu
85 90 95
Asp Ala Glu Lys Leu Ser Gln Leu Ser Ser Ser Leu Ile Ser Ser Phe 100 \hspace{1cm} 105 \hspace{1cm} 110
Ser
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 Leu Ile Arg Lys Pro Val Asp Phe Lys Lys Ile Lys Glu Arg Ile Arg 50 Asn His Lys Tyr Arg Ser Leu Asn Asp Leu Glu Lys Asp Val Met Leu 65 Cys Gln Asn Ala Gln Thr Phe Asn Leu Glu Gly Ser Leu Ile Tyr Glu Asp Ser Ile Val Leu Gln Ser Val Phe Thr Ser Val Arg Gln Lys 110 Cys Gln Lys

Ile Glu

<210> 38 <211> 113 <212> PRT

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<400> 38

Ser Pro Asn Pro Pro Lys Leu Thr Lys Gln Met Asn Ala Ile Ile Asp 1 10 15

Thr Val Ile Asn Tyr Lys Asp Ser Ser Gly Arg Gln Leu Ser Glu Val 20 25 30

Phe Ile Gln Leu Pro Ser Arg Lys Glu Leu Pro Glu Tyr Tyr Glu Leu 35 40 45

Ile Arg Lys Pro Val Asp Phe Lys Lys Ile Lys Glu Arg Ile Arg Asn 50 55 60

His Lys Tyr Arg Ser Leu Gly Asp Leu Glu Lys Asp Val Met Leu Leu 65 70 75 80

Cys His Asn Ala Gln Thr Phe Asn Leu Glu Gly Ser Gln Ile Tyr Glu 85 90 95

Asp Ser Ile Val Leu Gln Ser Val Phe Lys Ser Ala Arg Gln Lys Ile 100 105 110

Ala

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Ala Val Ile Lys Tyr Lys Asp Ser Ser Ser Gly Arg Gln Leu Ser Glu 20 25 30

Val Phe Ile Gln Leu Pro Ser Arg Lys Glu Leu Pro Glu Tyr Tyr Glu 35 40 45

Page 21

2459-1-003 SeqList 09-15-08.txt

Leu Ile Arg Lys Pro Val Asp Phe Lys Lys Ile Lys Glu Arg Ile Arg 50 Asn His Lys Tyr Arg Ser Leu Asn Asp Leu Glu Lys Asp Val Met Leu 65 Leu Cys Gln Asn Ala Gln Thr Phe Asn Leu Glu Val Ser Leu Ile Tyr 90 Glu Asp Ser Ile Val Leu Gln Ser Val Phe Thr Ser Val Arg Gln Lys 110 Lys

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Asp Ser Thr Phe Ser Leu Asp Gln Pro Gly Gly Thr Leu Asp Leu Thr 35 40 45

Leu Ile Arg Ala Arg Leu Gln Glu Lys Leu Ser Pro Pro Tyr Ser Ser 50 60

Pro Gln Glu Phe Ala Gln Asp Val Gly Arg Met Phe Lys Gln Phe Asn 70 75 80

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Arg Phe Phe Glu Thr Arg Met Asn Glu 100 105

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Ala Leu Phe Cys His Glu Pro Cys Arg Pro Leu His Gln Leu Ala Thr 20 25 30

Asp Ser Thr Phe Ser Met Glu Gln Pro Gly Gly Thr Leu Asp Leu Thr 35 40 45

Leu Ile Arg Ala Arg Leu Gln Glu Lys Leu Ser Pro Pro Tyr Ser Ser 50 60 Page 22

2459-1-003 SeqList 09-15-08.txt Pro Gln Glu Phe Ala Gln Asp Val Gly Arg Met Phe Lys Gln Phe Asn 65 70 75 80 Lys Leu Thr Glu Asp Lys Ala Asp Val Gln Ser Ile Ile Gly Leu Gln 85 90 95 Arg Phe Phe Glu Thr Arg Met Asn Asp 100 <210> 42 <211> <212> 108 PRT <213> Mus musculus <400> 42 Thr Lys Leu Thr Pro Ile Asp Lys Arg Lys Cys Glu Arg Leu Leu 1 10 15 Phe Leu Tyr Cys His Glu Met Ser Leu Ala Phe Gln Asp Pro Val Pro
20 25 30 Leu Thr Val Pro Asp Tyr Tyr Lys Ile Ile Lys Asn Pro Met Asp Leu 35 40 45 Ser Thr Ile Lys Lys Arg Leu Gln Glu Asp Tyr Cys Met Tyr Thr Lys 50 60 Pro Glu Asp Phe Val Ala Asp Phe Arg Leu Ile Phe Gln Asn Cys Ala 65 70 75 80 Glu Phe Asn Glu Pro Asp Ser Glu Val Ala Asn Ala Gly Ile Lys Leu 85 90 95 Glu Ser Tyr Phe Glu Glu Leu Leu Lys Asn Leu Tyr 100 105 <210> 43 18 <211> <212> PRT <213> Artificial Sequence <220> <223> synthetic bromodomain peptide <220> <221> xaa (1)..(1)<222> <223> Xaa can be any single amino acid <220> <221> Xaa <222> (2)..(2)Xaa can be any single amino acid

<222> (4)..(6) <223> Xaa is a maximum of three amino acids. Each of these can be any amino acid. Page 23

<220> <221>

Xaa

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2459-1-003 SeqList 09-15-08.txt
One may be missing.
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<222>
      (6)..(13)
<223> Xaa is a maximum of eight amino acids. Each of these can be any amino acid.
One, two, or three may be missing.
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      Xaa
<222>
       (7)..(7)
      Xaa is a single amino acid that can be Pro, Lys, or His.
<220>
<221>
      Xaa
<222>
      (8)..(8)
      Xaa is a single amino acid that can be any amino acid.
<220>
<221>
       Xaa
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       (10)..(10)
<223>
      Xaa is a single amino acid that can be a Tyr, Phe, or His.
<220>
<221> xaa
<222> (11)..(15)
<223> Xaa is five amino acids. Each of these can be any amino acid.
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<222> (13)..(13)
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<400> 43
Xaa Xaa Phe Xaa Pro Xaa Xaa Xaa Tyr Xaa Xaa Xaa Xaa Xaa Pro Xaa Asp
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       Artificial Sequence
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      synthetic bromodomain peptide
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Trp Pro Phe Met Glu Pro Val Lys Arg Thr Glu Ala Pro Gly Tyr Tyr
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1

Glu Val Ile Arg

20